

<110> Avicore Biotechnology Institute Inc.

<120> Recombinant ScFv Antibodies Specific to Eimeria spp. Responsible for Coccidiosis

<130> Avicore-USA-1

<150> KR 2001-52934

<151> 2001-08-30

<160> 40

<170> KopatentIn 1.71

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<212> DNA

<213> Artificial Sequence

<220>

<223> forward primer for PCR amplification of heavy chain variable region

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21

<210> 2

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<213> Artificial Sequence

<220>

<223> reverse primer for PCR amplification of heavy chain variable region

<400> 2  
gccgtgacgt tggacgagtc c

21

<210> 3  
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<400> 3  
taggacggtc agggttgtcc c

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<220>  
<223> reverse primer for PCR amplification of light chain variable region

<400> 4  
gcgcgtgactc agccgtcctc g

21

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<223> reverse primer for PCR amplification of heavy chain variable region

<400> 5  
ggcggagggtg gctctggcg tggcgatcg gccgtgacgt tggacgagtc c

51

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<400> 6  
ggaggagacg atgacttcgg t

21

<210> 7  
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<223> reverse primer for PCR amplification of light chain variable region

<400> 7  
gcgctgactc agccgtcctc g

21

<210> 8  
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<212> DNA  
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<220>  
<223> forward primer for PCR amplification of light chain variable region

<400> 8  
agagccacct ccgcctgaac cgccctccacc taggacggtc agggttgtcc c

51

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<220>  
<223> reverse primer for PCR amplification of heavy chain variable region

<400> 9  
gccgtgacgt tggacgagtc c

21

<210> 10  
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<400> 10  
agagccacct ccgcctgaac cgccctccacc ggaggagacg atgacttcgg t 51

<210> 11  
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<400> 11  
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<210> 12  
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<400> 12  
taggacggtc agggttgtcc c 21

<210> 13  
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<220>  
<223> reverse primer for PCR amplification of scFv

<400> 13  
gtcctcgcaa ctgcggccca gccgggcat ggccgcgtg actcagccgt cctcg 55

<210> 14  
<211> 39  
<212> DNA  
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<220>  
<223> forward primer for PCR amplification of scFv

<400> 14  
ggccacctt gcggccgcgg aggagacat gacttcggt 39

<210> 15  
<211> 55  
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<220>  
<223> reverse primer for PCR amplification of scFv

<400> 15  
gtcctcgcaa ctgcggccca gccgggcat ggccgccgtg acgttgacg agtcc 55

<210> 16  
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<212> DNA  
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<400> 16  
ggccacccttt gcggccgcta ggacggtcag gtttgtccc 39

<210> 17  
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<213> chicken hybridoma cell line 2-1

<220>  
<221> CDS  
<222> (1)..(369)

<400> 17  
gcc gtg acg ttg gac gag tcc ggg ggc ggc ctc cag acg ccc gga gga 48  
Ala Val Thr Leu Asp Glu Ser Gly Gly Leu Gln Thr Pro Gly Gly  
1 5 10 15

gcg ctc agc ctc gtc aag gcc tcc ggg ttc acc ttc agc agc cat 96  
Ala Leu Ser Leu Val Cys Lys Ala Ser Gly Phe Thr Phe Ser Ser His  
20 25 30

ggc atg atg tgg gtg cga cag acg ccc ggc aag ggg ctg gag tgg gtc				144
Gly Met Met Trp Val Arg Gln Thr Pro Gly Lys Glu Leu Trp Val				
35	40	45		
gcg ggt att agc aac act ggt act tac acg tac tac gcg ccg gcg gtg				192
Ala Gly Ile Ser Asn Thr Gly Thr Tyr Thr Tyr Ala Pro Ala Val				
50	55	60		
aag ggc cgt gcc acc atc tcg agg gac aac ggg cag agc aca gtg agg				240
Lys Gly Arg Ala Thr Ile Ser Arg Asp Asn Gly Gln Ser Thr Val Arg				
65	70	75	80	
ctg cag ctg aac aac ctc agg gct gag gac acc ggc acc tac tac tgc				288
Leu Gln Leu Asn Asn Leu Arg Ala Glu Asp Thr Gly Thr Tyr Tyr Cys				
85	90	95		
gcc aaa ggt ggt gct tat tgt gct ggt tgt ggt gac atc gac gca				336
Ala Lys Gly Gly Ala Tyr Cys Ala Gly Cys Gly Gly Asp Ile Asp Ala				
100	105	110		
tgg ggc cac ggg acc gaa gtc atc gtc tcc tcc				369
Trp Gly His Gly Thr Glu Val Ile Val Ser Ser				
115	120			
<210> 18				
<211> 123				
<212> PRT				
<213> chicken hybridoma cell line 2-1				
<400> 18				
Ala Val Thr Leu Asp Glu Ser Gly Gly Leu Gln Thr Pro Gly Gly				
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Ala Leu Ser Leu Val Cys Lys Ala Ser Gly Phe Thr Phe Ser Ser His				
20	25	30		

Gly Met Met Trp Val Arg Gln Thr Pro Gly Lys Gly Leu Glu Trp Val  
35 40 45

Ala Gly Ile Ser Asn Thr Gly Thr Tyr Thr Tyr Tyr Ala Pro Ala Val  
50 55 60

Lys Gly Arg Ala Thr Ile Ser Arg Asp Asn Gly Gln Ser Thr Val Arg  
65 70 75 80

Leu Gln Leu Asn Asn Leu Arg Ala Glu Asp Thr Gly Thr Tyr Tyr Cys  
85 90 95

Ala Lys Gly Gly Ala Tyr Cys Ala Gly Cys Gly Gly Asp Ile Asp Ala  
100 105 110

Trp Gly His Gly Thr Glu Val Ile Val Ser Ser  
115 120

<210> 19

<211> 372

<212> DNA

<213> chicken hybridoma cell line 5011

<220>

<221> CDS

<222> (1)..(372)

<400> 19

gcc gtg acg ttg gac gag tcc ggg ggc ggc ctc cag acg ccc gca gga  
Ala Val Thr Leu Asp Glu Ser Gly Gly Leu Gln Thr Pro Gly Gly  
1 5 10 15

gcg ctc agc ctc gtc tgc aag gcc tcc ggg ttc gac ttc agc agt tac  
Ala Leu Ser Leu Val Cys Lys Ala Ser Gly Phe Asp Phe Ser Ser Tyr  
96

20	25	30	
gac atg att tgg gtg cga cag gcg ccc ggc aag ggg ctg gaa tac gtc Asp Met Ile Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Tyr Val			144
35	40	45	
gcg ggt att aga agt gat ggt agt agc ata tac tac ggg gcg gcg gtg Ala Gly Ile Arg Ser Asp Gly Ser Ser Ile Tyr Tyr Gly Ala Ala Val			192
50	55	60	
aag ggc cgt gcc acc atc tcg agg gac aac ggg cag agc act ctg agg Lys Gly Arg Ala Thr Ile Ser Arg Asp Asn Gly Gln Ser Thr Leu Arg			240
65	70	75	80
ctg cag ctg aac aac ctc agg gct gag gac acc ggc acc tat tac tgc Leu Gln Leu Asn Asn Leu Arg Ala Glu Asp Thr Gly Thr Tyr Tyr Cys			288
85	90	95	
gcc aaa agt tct tat ggt agt tgg aga ggt tct act ggt gac atc gac Ala Lys Ser Ser Tyr Gly Ser Trp Arg Gly Ser Thr Gly Asp Ile Asp			336
100	105	110	
gca tgg ggc cac ggg acc gaa gtc atc gtc tcc tcc Ala Trp Gly His Gly Thr Glu Val Ile Val Ser Ser			372
115	120		
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<212> PRT			
<213> chicken hybridoma cell line 5D11			
<400> 20			
Ala Val Thr Leu Asp Glu Ser Gly Gly Gly Leu Gln Thr Pro Gly Gly			
1	5	10	15

Ala Leu Ser Leu Val Cys Lys Ala Ser Gly Phe Asp Phe Ser Ser Tyr  
20 25 30

Asp Met Ile Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Tyr Val  
35 40 45

Ala Gly Ile Arg Ser Asp Gly Ser Ser Ile Tyr Tyr Gly Ala Ala Val  
50 55 60

Lys Gly Arg Ala Thr Ile Ser Arg Asp Asn Gly Gln Ser Thr Leu Arg  
65 70 75 80

Leu Gln Leu Asn Asn Leu Arg Ala Glu Asp Thr Gly Thr Tyr Tyr Cys  
85 90 95

Ala Lys Ser Ser Tyr Gly Ser Trp Arg Gly Ser Thr Gly Asp Ile Asp  
100 105 110

Ala Trp Gly His Gly Thr Glu Val Ile Val Ser Ser  
115 120

<210> 21  
<211> 372  
<212> DNA  
<213> chicken hybridoma cell line 13C8

<220>  
<221> CDS  
<222> (1)..(372)

<400> 21  
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Ala Val Thr Leu Asp Glu Ser Gly Gly Gly Leu Gln Thr Pro Gly Gly  
1 5 10 15

ggg ctc agc ctc gtc tgc aag ggc tcc ggg ctc gac ttc agc agt tat			96
Gly Leu Ser Leu Val Cys Lys Gly Ser Gly Leu Asp Phe Ser Ser Tyr			
20	25	30	
gcc atg ggt tgg gtg cga cag gca ccc ggc aag ggg ctg gaa ttc gtc			144
Ala Met Gly Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Phe Val			
35	40	45	
gcg ggt att aaa aaa aat gat ggt agt tgg aca aac tac gcg ccg gcg			192
Ala Gly Ile Lys Lys Asn Asp Gly Ser Trp Thr Asn Tyr Ala Pro Ala			
50	55	60	
gtg cag ggc cgt gcc acc atc tcg agg gac aac ggg caa agc aca gtc			240
Val Gln Gly Arg Ala Thr Ile Ser Arg Asp Asn Gly Gln Ser Thr Val			
65	70	75	80
agg ctg cag ctg aac aac ctc agg gct gac gac acc ggc atc tac gtc			288
Arg Leu Gln Leu Asn Asn Leu Arg Ala Asp Asp Thr Gly Ile Tyr Val			
85	90	95	
tgc acc aga gat gtt aat agt ggt tac cct gat gct gct gac atc gac			336
Cys Thr Arg Asp Val Asn Ser Gly Tyr Pro Asp Ala Ala Asp Ile Asp			
100	105	110	
gca tgg ggc cac ggg acc gaa gtc atc gtc tcc tcc			372
Ala Trp Gly His Gly Thr Glu Val Ile Val Ser Ser			
115	120		
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<211> 124			
<212> PRT			
<213> chicken hybridoma cell line 13C8			
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1

5

10

15

Gly Leu Ser Leu Val Cys Lys Gly Ser Gly Leu Asp Phe Ser Ser Tyr

20

25

30

Ala Met Gly Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Phe Val

35

40

45

Ala Gly Ile Lys Lys Asn Asp Gly Ser Trp Thr Asn Tyr Ala Pro Ala

50

55

60

Val Gln Gly Arg Ala Thr Ile Ser Arg Asp Asn Gly Gln Ser Thr Val

65

70

75

80

Arg Leu Gln Leu Asn Asn Leu Arg Ala Asp Asp Thr Gly Ile Tyr Val

85

90

95

Cys Thr Arg Asp Val Asn Ser Gly Tyr Pro Asp Ala Ala Asp Ile Asp

100

105

110

Ala Trp Gly His Gly Thr Glu Val Ile Val Ser Ser

115

120

<210> 23

<211> 375

<212> DNA

<213> chicken hybridoma cell line 8C3

<220>

<221> CDS

<222> (1)..(375)

<400> 23

gcc gtg acg ttg gac gag tcc ggg ggc ggc ctc cag acg ccc gga gga

48

Ala Val Thr Leu Asp Glu Ser Gly Gly Leu Gln Thr Pro Gly Gly

1	5	10	15	
ggg ctc agc ctc gtc tgc aag gcc tcc ggg ttc tct atc ggc ggt tac Gly Leu Ser Leu Val Cys Lys Ala Ser Gly Phe Ser Ile Gly Gly Tyr				96
				20                    25                    30
				35                    40                    45
				144
atc atg cac tgg gtg cgc cag acg cct gga aag ggg ctg gaa tac gtt Ile Met His Trp Val Arg Gln Thr Pro Gly Lys Gly Leu Glu Tyr Val				192
				50                    55                    60
				192
gca ggt att gat gct ggt ggt agc aca tac tac ggg gcg gcg gtg Ala Gly Ile Asp Ala Gly Gly Ser Thr Tyr Tyr Gly Ala Ala Val				240
				65                    70                    75                    80
				240
cag ggc cgt gcc acc gtc tcg agg gac aac ggg cag agc aca ctg agg Gln Gly Arg Ala Thr Val Ser Arg Asp Asn Gly Gln Ser Thr Leu Arg				288
				85                    90                    95
				288
ctg cag ctg aac aac ctc agg ctg gag gac acc ggc acc tac ttc tgc Leu Gln Leu Asn Asn Leu Arg Leu Glu Asp Thr Gly Thr Tyr Phe Cys				336
				100                  105                  110
				336
gcc aaa gct tct cgg tgt ggc tat gat tgg tgt tct gct gat aac atc Ala Lys Ala Ser Arg Cys Gly Tyr Asp Trp Cys Ser Ala Asp Asn Ile				375
				115                  120                  125
				375

<210> 24

<211> 125

<212> PRT

<213> chicken hybridoma cell line 8C3

<400> 24

Ala Val Thr Leu Asp Glu Ser Gly Gly Gly Leu Gln Thr Pro Gly Gly

1 5 10 15

Gly Leu Ser Leu Val Cys Lys Ala Ser Gly Phe Ser Ile Gly Gly Tyr

20 25 30

Ile Met His Trp Val Arg Gln Thr Pro Gly Lys Gly Leu Glu Tyr Val

35 40 45

Ala Gly Ile Asp Ala Gly Gly Ser Thr Tyr Tyr Gly Ala Ala Val

50 55 60

Gln Gly Arg Ala Thr Val Ser Arg Asp Asn Gly Gln Ser Thr Leu Arg

65 70 75 80

Leu Gln Leu Asn Asn Leu Arg Leu Glu Asp Thr Gly Thr Tyr Phe Cys

85 90 95

Ala Lys Ala Ser Arg Cys Gly Tyr Asp Trp Cys Ser Ala Asp Asn Ile

100 105 110

Asp Ala Trp Gly His Gly Thr Glu Val Ile Val Ser Ser

115 120 125

<210> 25

<211> 324

<212> DNA

<213> chicken hybridoma cell line 2-1

<220>

<221> CDS

<222> (1)..(324)

<400> 25

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Ala	Leu	Thr	Gln	Pro	Ser	Ser	Val	Ser	Ala	Asn	Pro	Gly	Glu	Thr	Val		
1		5					10								15		
aag	atc	acc	tgc	tcc	ggg	ggt	ggc	agc	tac	gct	gga	agt	tac	tat	tat		96
Lys	Ile	Thr	Cys	Ser	Gly	Gly	Gly	Ser	Tyr	Ala	Gly	Ser	Tyr	Tyr	Tyr		
		20					25								30		
ggc	tgg	tac	cag	cag	aag	gca	cct	gcc	agt	gcc	cct	gtc	act	gtg	atc		144
Gly	Trp	Tyr	Gln	Gln	Lys	Ala	Pro	Ala	Ser	Ala	Pro	Val	Thr	Val	Ile		
		35				40									45		
tat	gac	aac	acc	aac	aga	ccc	tcg	aac	atc	cct	tca	cga	ttc	tcc	ggt		192
Tyr	Asp	Asn	Thr	Asn	Arg	Pro	Ser	Asn	Ile	Pro	Ser	Arg	Phe	Ser	Gly		
		50				55									60		
tcc	cta	tcc	ggc	tcc	aca	aac	aca	tta	acc	atc	act	ggg	gtc	caa	gtc		240
Ser	Leu	Ser	Gly	Ser	Thr	Asn	Thr	Leu	Thr	Ile	Thr	Gly	Val	Gln	Val		
		65			70			75							80		
gag	gac	gag	gct	tat	tac	tgt	ggg	agg	ttc	gac	agg	agt	tat	gtt		288	
Glu	Asp	Glu	Ala	Val	Tyr	Tyr	Cys	Gly	Ser	Phe	Asp	Ser	Ser	Tyr	Val		
		85				90									95		
ggt	ata	ctt	ggg	gcc	ggg	aca	acc	ctg	acc	gtc	cta						324
Gly	Ile	Leu	Gly	Ala	Gly	Thr	Thr	Leu	Thr	Val	Leu						
		100				105											

<210> 26

<211> 108

<212> PRT

<213> chicken hybridoma cell line 2-1

<400> 26

Ala Leu Thr Gln Pro Ser Ser Val Ser Ala Asn Pro Gly Glu Thr Val

1	5	10	15	
Lys Ile Thr Cys Ser Gly Gly Ser Tyr Ala Gly Ser Tyr Tyr Tyr				
	20	25	30	
Gly Trp Tyr Gln Gln Lys Ala Pro Ala Ser Ala Pro Val Thr Val Ile				
	35	40	45	
Tyr Asp Asn Thr Asn Arg Pro Ser Asn Ile Pro Ser Arg Phe Ser Gly				
	50	55	60	
Ser Leu Ser Gly Ser Thr Asn Thr Leu Thr Ile Thr Gly Val Gln Val				
	65	70	75	80
Glu Asp Glu Ala Val Tyr Tyr Cys Gly Ser Phe Asp Ser Ser Tyr Val				
	85	90	95	
Gly Ile Leu Gly Ala Gly Thr Thr Leu Thr Val Leu				
	100	105		

<210> 27  
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<212> DNA  
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<220>  
<221> CDS  
<222> (1)...(312)

<400> 27  
gcg ctg act cag ccg tcc tcg gtg tca gca aac ctg gga gaa acc gtc 48  
Ala Leu Thr Gln Pro Ser Ser Val Ser Ala Asn Leu Gly Glu Thr Val  
1 5 10 15  
gaa atc acc tgc tcc ggg ggc agg tat agg tat ggc tgg tat cag cag 96

Glu	Ile	Thr	Cys	Ser	Gly	Gly	Arg	Tyr	Arg	Tyr	Gly	Trp	Tyr	Gln	Gln	
20																
aag	tca	tct	ggc	agt	gcc	cct	gtc	act	gtg	atc	tat	gac	aac	gac	aag	144
Lys	Ser	Ser	Gly	Ser	Ala	Pro	Val	Thr	Val	Ile	Tyr	Asp	Asn	Asp	Lys	
35																
aga	ccc	tcg	gac	atc	cct	tca	cga	ttc	tcc	ggt	tcc	aaa	tcc	gac	tcc	192
Arg	Pro	Ser	Asp	Ile	Pro	Ser	Arg	Phe	Ser	Gly	Ser	Lys	Ser	Asp	Ser	
50																
acg	ggc	aca	tta	acc	atc	act	ggg	gtc	caa	gcc	gag	gac	gag	gct	gtc	240
Thr	Gly	Thr	Leu	Thr	Ile	Thr	Gly	Val	Gln	Ala	Glu	Asp	Glu	Ala	Val	
65																
tat	tac	tgt	ggg	aat	gca	gac	aac	aat	act	tac	gat	cct	ata	ttt	ggg	288
Tyr	Tyr	Cys	Gly	Asn	Ala	Asp	Asn	Asn	Thr	Tyr	Asp	Pro	Ile	Phe	Gly	
85																
gcc	ggg	aca	acc	ctg	acc	gtc	cta									312
Ala	Gly	Thr	Thr	Leu	Thr	Val	Leu									
100																
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<211> 104																
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<213> chicken hybridoma cell line 5D11																
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Ala Leu Thr Gln Pro Ser Ser Val Ser Ala Asn Leu Gly Glu Thr Val																
1		5													15	
Glu Ile Thr Cys Ser Gly Gly Arg Tyr Arg Tyr Gly Trp Tyr Gln Gln																
20															30	

Lys Ser Ser Gly Ser Ala Pro Val Thr Val Ile Tyr Asp Asn Asp Lys  
35 40 45

Arg Pro Ser Asp Ile Pro Ser Arg Phe Ser Gly Ser Lys Ser Asp Ser  
50 55 60

Thr Gly Thr Leu Thr Ile Thr Gly Val Gln Ala Glu Asp Glu Ala Val  
65 70 75 80

Tyr Tyr Cys Gly Asn Ala Asp Asn Asn Thr Tyr Asp Pro Ile Phe Gly  
85 90 95

Ala Gly Thr Thr Leu Thr Val Leu  
100

<210> 29

<211> 324

<212> DNA

<213> chicken hybridoma cell line 13C8

<220>

<221> CDS

<222> (1)..(324)

<400> 29

gcg ctg act cag ccg tcc tcg gtg tca gca aac ctg gga gga acc gtc 48  
Ala Leu Thr Gln Pro Ser Ser Val Ser Ala Asn Leu Gly Gly Thr Val  
1 5 10 15

aag atc acc tgc tcc ggg ggc agc tat ggc tat ggc tgg ttc cag cag 96  
Lys Ile Thr Cys Ser Gly Gly Ser Tyr Gly Tyr Gly Trp Phe Gln Gln  
20 25 30

aag tca cct ggc agt gcc cct gtc cct gtg atc tac tgg aac aac aag 144  
Lys Ser Pro Gly Ser Ala Pro Val Pro Val Ile Tyr Trp Asn Asn Lys

35

40

45

aga ccc tcg gac atc cct tca cga ttc tcc ggt tcc aaa tcc ggc tcc 192

Arg Pro Ser Asp Ile Pro Ser Arg Phe Ser Gly Ser Lys Ser Gly Ser

50

55

60

aca gcc aca tta acc atc act ggg gtc cga gcc gag gac gag gct gtc 240

Thr Ala Thr Leu Thr Ile Thr Gly Val Arg Ala Glu Asp Glu Ala Val

65

70

75

80

tat tac tgt ggg aat gca gac agc aat act gct gat agt gat tat gtt 288

Tyr Tyr Cys Gly Asn Ala Asp Ser Asn Thr Ala Asp Ser Asp Tyr Val

85

90

95

ggt ata ttt ggg gcc ggg aca acc ctg acc gtc cta 324

Gly Ile Phe Gly Ala Gly Thr Thr Leu Thr Val Leu

100

105

<210> 30

<211> 108

<212> PRT

<213> chicken hybridoma cell line 13C8

<400> 30

Ala Leu Thr Gln Pro Ser Ser Val Ser Ala Asn Leu Gly Gly Thr Val

1

5

10

15

Lys Ile Thr Cys Ser Gly Gly Ser Tyr Gly Tyr Gly Trp Phe Gln Gln

20

25

30

Lys Ser Pro Gly Ser Ala Pro Val Pro Val Ile Tyr Trp Asn Asn Lys

35

40

45

Arg Pro Ser Asp Ile Pro Ser Arg Phe Ser Gly Ser Lys Ser Gly Ser

50

55

60

Thr Ala Thr Leu Thr Ile Thr Gly Val Arg Ala Glu Asp Glu Ala Val  
65 70 75 80

Tyr Tyr Cys Gly Asn Ala Asp Ser Asn Thr Ala Asp Ser Asp Tyr Val  
85 90 95

Gly Ile Phe Gly Ala Gly Thr Thr Leu Thr Val Leu  
100 105

<210> 31

<211> 315

<212> DNA

<213> chicken hybridoma cell line 8C3

<220>

<221> CDS

<222> (1)..(315)

<400> 31

gcg ctg act caa ccg tcc tcg gtg tca gcg atc ccg gga gaa acc gtc 48  
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1 5 10 15

gag atc acc tgc tcc ggg ggt aac aac tac tat ggc tgg tat cag cag 96  
Glu Ile Thr Cys Ser Gly Gly Asn Asn Tyr Tyr Gly Trp Tyr Gln Gln  
20 25 30

aaa tca cct ggc agt gcc cct gtc act gtg atc tac tac aac aac aag 144  
Lys Ser Pro Gly Ser Ala Pro Val Thr Val Ile Tyr Tyr Asn Asn Lys  
35 40 45

aga ccc tcg gac atc cct tca cga ttc tcc ggt tcc aaa ccc ggc tcc 192  
Arg Pro Ser Asp Ile Pro Ser Arg Phe Ser Gly Ser Lys Pro Gly Ser  
50 55 60

aca aac aca tta acc atc act ggg gtc cga gcc gag gac gag gct gtc 240  
Thr Asn Thr Leu Thr Ile Thr Gly Val Arg Ala Glu Asp Glu Ala Val  
65 70 75 80

tat ttc tgt ggt gcc tgg gaa agt agt cct att tat gtt ggt ata ttt 288  
Tyr Phe Cys Gly Ala Trp Glu Ser Ser Pro Ile Tyr Val Gly Ile Phe  
85 90 95

ggg gcc ggg aca acc ctg acc gtc cta 315  
Gly Ala Gly Thr Thr Leu Thr Val Leu  
100 105

<210> 32  
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<212> PRT  
<213> chicken hybridoma cell line 8C3

<400> 32  
Ala Leu Thr Gln Pro Ser Ser Val Ser Ala Ile Pro Gly Glu Thr Val  
1 5 10 15

Glu Ile Thr Cys Ser Gly Gly Asn Asn Tyr Tyr Gly Trp Tyr Gln Gln  
20 25 30

Lys Ser Pro Gly Ser Ala Pro Val Thr Val Ile Tyr Tyr Asn Asn Lys  
35 40 45

Arg Pro Ser Asp Ile Pro Ser Arg Phe Ser Gly Ser Lys Pro Gly Ser  
50 55 60

Thr Asn Thr Leu Thr Ile Thr Gly Val Arg Ala Glu Asp Glu Ala Val  
65 70 75 80

Tyr Phe Cys Gly Ala Trp Glu Ser Ser Pro Ile Tyr Val Gly Ile Phe

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Gly Ala Gly Thr Thr Leu Thr Val Leu

100

105

<210> 33

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> reverse primer for PCR amplification of heavy chain variable  
region

<400> 33

Ala Val Thr Leu Asp Glu Ser

1

5

<210> 34

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> forward primer for PCR amplification of heavy chain variable  
region

<400> 34

Ser Ser Val Ile Val Glu Thr

1

5

<210> 35  
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<220>  
<223> reverse primer for PCR amplification of light chain variable region

<400> 35  
Ala Leu Thr Gln Pro Ser Ser  
1 5

<210> 36  
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<220>  
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<400> 36  
Leu Val Thr Leu Thr Thr Gly  
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<400> 37

gcc gtg acg ttg gac gag tcc ggg ggc ggc ctc cag acg ccc gga aga  
Ala Val Thr Leu Asp Glu Ser Gly Gly Gly Leu Gln Thr Pro Gly Arg  
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48

gcg ctc agc ctc gtc tgc aag gcc tcc ggg ttc acc ttc agc agt tat  
Ala Leu Ser Leu Val Cys Lys Ala Ser Gly Phe Thr Phe Ser Ser Tyr  
20 25 30

96

ggc atg gtc tgg gtc cga cag gcg ccc ggc aag ggg ctg gaa tac gtc  
Gly Met Val Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Tyr Val  
35 40 45

144

gct gaa att atc aca act ggt aga gac aca tgg tat ggg acg gcg gtc  
Ala Glu Ile Ile Thr Thr Gly Arg Asp Thr Trp Tyr Gly Thr Ala Val  
50 55 60

192

aag ggc cgt gcc acc atc tcg agg gac aac ggg cag agt aca gtc agg  
Lys Gly Arg Ala Thr Ile Ser Arg Asp Asn Gly Gln Ser Thr Val Arg  
65 70 75 80

240

ctg cag ctg aac aac ctc agg gct gaa gac acc ggc atc tac tac tgc  
Leu Gln Leu Asn Asn Leu Arg Ala Glu Asp Thr Gly Ile Tyr Tyr Cys  
85 90 95

288

gcc aaa tgc agt tat gag tgt act agt agt tgt tgg ggt tat act gat  
Ala Lys Cys Ser Tyr Glu Cys Thr Ser Ser Cys Trp Gly Tyr Thr Asp  
100 105 110

336

atg atc gac gca tgg ggc cac ggg acc gaa gtc atc gtc tcc tcc  
Met Ile Asp Ala Trp Gly His Gly Thr Glu Val Ile Val Ser Ser  
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381

<210> 38

<211> 127

<212> PRT

<213> chicken hybridoma cell line 60-12-G10

<400> 38

Ala Val Thr Leu Asp Glu Ser Gly Gly Gly Leu Gln Thr Pro Gly Arg

1 5 10 15

Ala Leu Ser Leu Val Cys Lys Ala Ser Gly Phe Thr Phe Ser Ser Tyr

20 25 30

Gly Met Val Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Tyr Val

35 40 45

Ala Glu Ile Ile Thr Thr Gly Arg Asp Thr Trp Tyr Gly Thr Ala Val

50 55 60

Lys Gly Arg Ala Thr Ile Ser Arg Asp Asn Gly Gln Ser Thr Val Arg

65 70 75 80

Leu Gln Leu Asn Asn Leu Arg Ala Glu Asp Thr Gly Ile Tyr Tyr Cys

85 90 95

Ala Lys Cys Ser Tyr Glu Cys Thr Ser Ser Cys Trp Gly Tyr Thr Asp

100 105 110

Met Ile Asp Ala Trp Gly His Gly Thr Glu Val Ile Val Ser Ser

115 120 125

<210> 39

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<213> chicken hybridoma cell line 6D-12-G10

<220>

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<222> (1)...(312)

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Ala Leu Thr Gln Pro Ser Ser Val Ser Ala Asn Leu Gly Gly Thr Val

48

1                    5                    10                    15

aag atc acc tgc tcc ggg agt agt ggc agc tat ggc tgg tat cag cag		
Lys Ile Thr Cys Ser Gly Ser Ser Gly Ser Tyr Gly Trp Tyr Gln Gln		
20	25	30

96

144

192

210

tat ttc tgt gag agt aca gac tac agt agt act gat ata ttt ggg gcc  
 Tyr Phe Cys Glu Ser Thr Asp Tyr Ser Ser Thr Asp Ile Phe Gly Ala  
           85                  90                  95

288

ggg aca acc ctg acc gtc cta ggt  
 Gly Thr Thr Leu Thr Val Leu Gly

310

<210> 40

<211> 104

<212> PRT

<213> chicken hybridoma cell line 6D-12-G10

<400> 40

Ala Leu Thr Gln Pro Ser Ser Val Ser Ala Asn Leu Gly Gly Thr Val

1 5 10 15

Lys Ile Thr Cys Ser Gly Ser Ser Gly Ser Tyr Gly Trp Tyr Gln Gln

20 25 30

Lys Ser Pro Gly Ser Ala Pro Val Thr Val Ile Tyr Tyr Asn Asp Lys

35 40 45

Arg Pro Ser Asp Ile Pro Ser Arg Phe Ser Gly Ser Lys Ser Gly Ser

50 55 60

Thr Gly Thr Leu Thr Ile Thr Gly Val Gln Ala Glu Asp Glu Ala Val

65 70 75 80

Tyr Phe Cys Glu Ser Thr Asp Tyr Ser Ser Thr Asp Ile Phe Gly Ala

85 90 95

Gly Thr Thr Leu Thr Val Leu Gly

100